



SEQUENCE LISTING

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Pollitt, N. Stephen

<120> VASCULAR ENDOTHELIAL GROWTH FACTOR DIMERS

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<141> 2003-12-31

<150> 09/575,199

<151> 2000-05-18

<150> US 60/177,407

<151> 2000-01-20

<150> US 60/135,312

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1 5 10 15	
tac ctc cac cat gcc aag tgg tcc cag gct gca ccc atg gca gaa gga	96
Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly	
20 25 30	
gga ggg cag aat cat cac gaa gtg gtg aag ttc atg gat gtc tat cag	144
Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln	
35 40 45	
cgc agc tac tgc cat cca atc gag acc ctg gtg gac atc ttc cag gag	192
Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu	
50 55 60	
tac cct gat gag atc gag tac atc ttc aag cca tcc tgt gtg ccc ctg	240
Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu	
65 70 75 80	

atg cga tgc ggg ggc tgc tgc aat gac gag ggc ctg gag tgt gtg ccc	288
Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro	
85 90 95	

act gag gag tcc aac atc acc atg cag att atg cgg atc aaa cct cac	336
Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His	
100 105 110	

caa ggc cag cac ata gga gag atg agc ttc cta cag cac aac aaa tgt	384
Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys	
115 120 125	

gaa tgc aga cca aag aaa gat aga gca aga caa gaa aaa tgt gac aag	432
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ccg agg cgg tga	444
Pro Arg Arg	
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Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly	
20 25 30	
Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln	
35 40 45	
Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu	
50 55 60	
Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu	
65 70 75 80	
Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro	
85 90 95	
Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Arg Ile Lys Pro His	
100 105 110	
Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys	
115 120 125	
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<220>
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ttc atg gat gtc tat cag cgc agc tac tgc cat cca atc gag acc ctg	96
Phe Met Asp Val Tyr Gln Arg Ser Tyr Cys His Pro Ile Glu Thr Leu	
20 25 30	
gtg gac atc ttc cag gag tac cct gat gag atc gag tac atc ttc aag	144
Val Asp Ile Phe Gln Glu Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys	
35 40 45	
cca tcc tgt gtg ccc ctg atg cga tgc ggg ggc tgc tgc aat gac gag	192
Pro Ser Cys Val Pro Leu Met Arg Cys Gly Gly Cys Cys Asn Asp Glu	
50 55 60	
ggc ctg gag tgt gtg ccc act gag gag tcc aac atc acc atg cag att	240
Gly Leu Glu Cys Val Pro Thr Glu Glu Ser Asn Ile Thr Met Gln Ile	
65 70 75 80	
atg cgg atc aaa cct cac caa ggc cag cac ata gga gag atg agc ttc	288
Met Arg Ile Lys Pro His Gln Gly Gln His Ile Gly Glu Met Ser Phe	
85 90 95	
cta cag cac aac aaa tgt gaa tgc aga cca aag aaa gat aga gca aga	336
Leu Gln His Asn Lys Cys Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg	
100 105 110	
caa gaa aaa tgt gac aag ccg agg cgg tga	366
Gln Glu Lys Cys Asp Lys Pro Arg Arg	
115 120	